



# SEQUENCE LISTING

<110> Williams, John G. K.  
LI-COR, Inc.

<120> Composition and Method for Nucleic Acid Sequencing

<130> 020031-003110US

<140> US 10/821,689

<141> 2004-04-08

<150> US 60/461,522

<151> 2003-04-08

<150> US 60/462,988

<151> 2003-04-14

<160> 23

<170> PatentIn Ver. 2.1

<210> 1

<211> 89

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:synthetic target  
nucleic acid, single molecule in microtiter plate well

<400> 1

tatgaaaatt ttccggttta aggcgtttcc gttcttcttc gtcataactt aatgttttta 60  
tttaaaatac cctctgaaaa gaaaggaaa 89

<210> 2

<211> 89

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:synthetic target  
nucleic acid, single molecule in microtiter plate well

<400> 2

cgacaggtgc tgaaagcgag gctttttggc ctctgtcggt tcctttctct gtttttgtcc 60  
gtggaatgaa caatggaagt caacaaaaa 89

<210> 3

<211> 89

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:synthetic target  
nucleic acid, single molecule in microtiter plate well

<400> 3  
gcagctggct gacattttcg gtgcgagtat ccgtaccatt cagaactggc aggaacaggg 60  
aatgcccgtt ctgcgaggcg gtggcaagg 89

<210> 4  
<211> 89  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic target  
nucleic acid, single molecule in microtiter plate well

<400> 4  
gtaatgaggt gctttatgac tctgccgccg tcataaaatg gtatgccgaa agggatgctg 60  
aaattgagaa cgaaaagctg cgccgggag 89

<210> 5  
<211> 21  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic amino  
acid anchor sequence

<220>  
<221> MOD\_RES  
<222> (11)  
<223> Xaa = p-acetyl-L-phenylalanine (pa-Phe)

<400> 5  
Leu Leu Ser Lys Lys Arg Ser Leu Cys Cys Xaa Cys Thr Val Ile Val  
1 5 10 15  
Tyr Val Thr Asp Thr  
20

<210> 6  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:first  
double-stranded oligonucleotide adaptor

<220>  
<221> modified\_base  
<222> (1)  
<223> n = biotinylated c

<400> 6  
ngccacatta cacttcctaa cacgt 25

<210> 7  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:complement of  
       first double-stranded oligonucleotide adaptor  
  
 <400> 7  
 cgtgtagga agtgtaatgt ggcg 24  
  
 <210> 8  
 <211> 25  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:second  
       double-stranded oligonucleotide adaptor  
  
 <400> 8  
 cagtaggtag tcaaggctag agtct 25  
  
 <210> 9  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:complement of  
       second double-stranded oligonucleotide adaptor  
  
 <400> 9  
 gactctagcc ttgactacct actg 24  
  
 <210> 10  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:ligated DNA  
       product  
  
 <220>  
 <221> modified\_base  
 <222> (1)  
 <223> n = biotinylated c  
  
 <220>  
 <221> modified\_base  
 <222> (26)..(30)  
 <223> n = g, a, c or t  
  
 <400> 10  
 ngccacatta cacttcctaa cacgtnnnnn 30

<210> 11  
 <211> 33  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:ligated DNA  
         product  
  
 <220>  
 <221> modified\_base  
 <222> (1)..(5)  
 <223> n = g, a, c or t  
  
 <400> 11  
 nnnnnagact ctagccttga ctacctactg aaa 33

<210> 12  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:ligated DNA  
         product, unbiotinylated DNA strand eluted at  
         alkaline pH  
  
 <220>  
 <221> modified\_base  
 <222> (1)..(5)  
 <223> n = g, a, c or t  
  
 <400> 12  
 nnnnnacgtg ttaggaagtg taatgtggcg 30

<210> 13  
 <211> 30  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:ligated DNA  
         product, unbiotinylated DNA strand eluted at  
         alkaline pH  
  
 <220>  
 <221> modified\_base  
 <222> (1)  
 <223> n = 5' phosphorylated c  
  
 <220>  
 <221> modified\_base  
 <222> (26)..(30)  
 <223> n = g, a, c or t  
  
 <400> 13  
 nagtaggtag tcaaggctag agtctnnnnn 30

<210> 14  
 <211> 59  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:primed circular  
         template strand, eluted strands circularized  
  
 <220>  
 <221> modified\_base  
 <222> (1)..(59)  
 <223> n = g, a, c or t  
  
 <400> 14  
 nnnnncgtgt taggaagtgt aatgtggcgc agtaggtagt caaggctaga gtctnnnnn 59  
  
 <210> 15  
 <211> 49  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:primer oligo  
         complementary to both adaptors  
  
 <400> 15  
 agactctagc cttgactacc tactgcgccca cattacactt cctaacacg 49  
  
 <210> 16  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:T7 DNA  
         polymerase gene forward amplification primer  
         encoding exonuclease mutations  
  
 <400> 16  
 atgatcgttt ctgccatcgc agctaac 27  
  
 <210> 17  
 <211> 16  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence:T7 DNA  
         polymerase gene reverse amplification primer  
  
 <400> 17  
 tcagtggcaa atcgcc 16

<210> 18  
<211> 75  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic  
oligonucleotide encoding Strep-Tag II sequence  
overlapping 5'-end N-terminus of amplified T7  
polymerase gene with 2 exo- mutations

<220>  
<221> CDS  
<222> (1)..(75)  
<223> Strep-Tag II peptide, spacer and T7 polymerase  
N-terminus overlap with 2 exo- mutations

<400> 18  
atg tcc aac tgg tcc cac ccg cag ttc gaa aaa ggt gga ggt tcc gct 48  
Met Ser Asn Trp Ser His Pro Gln Phe Glu Lys Gly Gly Gly Ser Ala  
1 5 10 15  
  
atg atc gtt tct gcc atc gca gct aac 75  
Met Ile Val Ser Ala Ile Ala Ala Asn  
20 25

<210> 19  
<211> 25  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Strep-Tag II peptide, spacer and T7 polymerase  
N-terminus overlap with 2 exo- mutations

<400> 19  
Met Ser Asn Trp Ser His Pro Gln Phe Glu Lys Gly Gly Gly Ser Ala  
1 5 10 15  
  
Met Ile Val Ser Ala Ile Ala Ala Asn  
20 25

<210> 20  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:overlapping PCR  
synthetic oligonucleotide StrepTag forward primer

<400> 20  
atgtccaact ggtcccaccc 20

<210> 21  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic  
oligonucleotide sample primer derived from cystic  
fibrosis transmembrane conductance regulator gene

<400> 21  
tactataaaa gaaattacca c

21

<210> 22  
<211> 22  
<212> RNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic  
oligonucleotide sample template derived from  
cystic fibrosis transmembrane conductance  
regulator gene normal allele

<400> 22  
gugguaauuu cuuuuauagu ag

22

<210> 23  
<211> 22  
<212> RNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:synthetic  
oligonucleotide sample template derived from  
cystic fibrosis transmembrane conductance  
regulator gene (delta)F508 deletion mutant

<400> 23  
gugguaauuu cuuuuauagu aa

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